

Sequence Revision History

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Revision history for "U24231"				
GI	Version	Update Date	Status	
809486	1	Feb 2 1996 1:09	Live	
809486	1	May 20 1995 2:17	Dead	

Accession <u>U24231</u> was first seen at <u>NCBI</u> on May 20 1995 2:17

Entrez Nucleotide

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Batch Entrez: Upload a file of GI or accession numbers to retrieve sequences

Check sequence revision history

How to create WWW links to Entrez

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Reference sequence project

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Revised: July 5, 2002.

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Revision history for "Q13158"				
GI	Version	Update Date	Status	
2498355	N/A	Oct 1 2002 9:49	Live	
2498355	N/A	Apr 1 2002 17:54	Dead	
2498355	N/A	Jun 25 2001 15:20	Dead	
2498355	N/A	Apr 16 2001 14:46	Dead	
2498355	N/A	Feb 1 2001 12:01	Dead	
2498355	N/A	Nov 1 1999 13:03	Dead	
2498355	N/A	Jul 21 1998 5:35	Dead	
2498355	N/A	Jun 4 1998 4:24	Dead	
2498355	N/A	Feb 5 1998 14:29	Dead	
2498355	N/A	Oct 8 1997 19:11	Dead	

SeqId was first seen at NCBI on Oct 8 1997 19:11

Revised: August 5, 2002.

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□1: Cell 1995 May 19;81(4):505-12

Related Articles, Links

FULL @ CELL

FADD, a novel death domain-containing protein, interacts with the death domain of Fas and initiates apoptosis.

Chinnaiyan AM, O'Rourke K, Tewari M, Dixit VM.

Department of Pathology, University of Michigan Medical School, Ann Arbor 48109, USA.

Using the cytoplasmic domain of Fas in the yeast two-hybrid system, we have identified a novel interacting protein, FADD, which binds Fas and Fas-FD5, a mutant of Fas possessing enhanced killing activity, but not the functionally inactive mutants Fas-LPR and Fas-FD8. FADD contains a death domain homologous to the death domains of Fas and TNFR-1. A point mutation in FADD, analogous to the lpr mutation of Fas, abolishes its ability to bind Fas, suggesting a death domain to death domain interaction. Overexpression of FADD in MCF7 and BJAB cells induces apoptosis, which, like Fas-induced apoptosis, is blocked by CrmA, a specific inhibitor of the interleukin-1 beta-converting enzyme. These findings suggest that FADD may play an important role in the proximal signal transduction of Fas.

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MeSH Terms:

- Amino Acid Sequence
- Antigens, CD95
- Antigens, Surface/metabolism*
- Antigens, Surface/genetics
- Apoptosis*
- Base Sequence
- Binding Sites/genetics
- Carrier Proteins/metabolism
- Carrier Proteins/isolation & purification*
- Carrier Proteins/genetics
- Cell Line
- · Cloning, Molecular
- Human
- Molecular Sequence Data
- Point Mutation

- Saccharomyces cerevisiae
- Serpins/pharmacology
- Signal Transduction
- · Support, Non-U.S. Gov't

Substances:

- interleukin-1beta-converting enzyme inhibitor
- Serpins
- MORT1 protein
- · Carrier Proteins
- · Antigens, Surface
- Antigens, CD95

Secondary source id:

GENBANK/U24231

PMID: 7538907 [PubMed - indexed for MEDLINE]



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Feb 19 2003 14:38:47